APPENDIX A

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RESULT 2
US-09-946-374-85
; Sequence 85, Application US/09946374
; Publication No. US20030073129A1
; GENERAL INFORMATION:
;  APPLICANT: Baker, Kevin P.
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Goddard, Audrey
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth J.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Smith, Victoria
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Watanabe, Colin K.
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: P2830P1C1
;  CURRENT APPLICATION NUMBER: US/09/946,374
;  CURRENT FILING DATE:  2001-09-04
;  PRIOR APPLICATION NUMBER: 60/098716
;  PRIOR FILING DATE: 1998-09-01
;  PRIOR APPLICATION NUMBER: 60/098723
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;   PRIOR FILING DATE: 1998-10-26
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;   PRIOR APPLICATION NUMBER: 60/105807

  Query Match          98.3%;  Score 3261;  DB 10;  Length 3316;
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  Matches 3311;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

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Qy    1981  TTAGTAGGAGTGGTGGTTCCTGACACAGATGTACTTCCCTCATTTGCAGCCAAGCTTGGG  2040
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Qy    2161  TTTCTTCATCCAGAGCCATTTTCCATTGAAAATGGGCTCTTGACACCAACATTGAAAGCA  2220
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Qy    2221  AAGCGAGGAGAGCTTTCCAAATACTTTCGGACCCAAATTGACAGCCTGTATGAGCACATC  2280
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Qy    2521  ACTTTTATCAACATGCCTGTCTTCAAGATCCCAGTTTATGTTCTGTGTCCTTCCTCATGA  2580
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Db    2525  ACTTTTATCAACATGCCTGTCTTCAAGATCCCAGTTTATGTTCTGTGTCCTTCCTCATGA  2584

Qy    2581  TTTCCAACCTTAATACTATTAGTAACCACAAGTTCAAGGGTCAAAGGGACCCTCTGTGCC  2640
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Db    2645  TTCTTCTTTGTTTTGTGATAAACATAACTTGCCAACAGTCTCTATGCTTATTTACATCTT  2704

Qy    2701  CTACTGTTCAAACTAAGAGATTTTTAAATTCTGAAAAACTGCTTACAATTCATGTTTTCT  2760
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Qy    2761  AGCCACTCCACAAACCACTAAAATTTTAGTTTTAGCCTATCACTCATGTCAATCATATCT  2820
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Qy    2821  ATGAGACAAATGTCTCCGATGCTCTTCTGCGTAAATTAAATTGTGTACTGAAGGGAAAAG  2880
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Db    2825  ATGAGACAAATGTCTCCGATGCTCTTCTGCGTAAATTAAATTGTGTACTGAAGGGAAAAG  2884
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Qy    2881 TTTGATCATACCAAACATTTCCTAAACTCTCTAGTTAGATATCTGACTTGGGAGTATTAA 2940
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2885 TTTGATCATACCAAACATTTCCTAAACTCTCTAGTTAGATATCTGACTTGGGAGTATTAA 2944

Qy    2941 AAATTGGGTCTATGACATACTGTCCAAAAGGAATGCTGTTCTTAAAGCATTATTTACAGT 3000
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2945 AAATTGGGTCTATGACATACTGTCCAAAAGGAATGCTGTTCTTAAAGCATTATTTACAGT 3004

Qy    3001 AGGAACTGGGGAGTAAATCTGTTCCCTACAGTTTGCTGCTGAGCTGGAAGCTGTGGGGGA 3060
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3005 AGGAACTGGGGAGTAAATCTGTTCCCTACAGTTTGCTGCTGAGCTGGAAGCTGTGGGGGA 3064

Qy    3061 AGGAGTTGACAGGTGGGCCCAGTGAACTTTTCCAGTAAATGAAGCAAGCACTGAATAAAA 3120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3065 AGGAGTTGACAGGTGGGCCCAGTGAACTTTTCCAGTAAATGAAGCAAGCACTGAATAAAA 3124

Qy    3121 ACCTCCTGAACTGGGAACAAAGATCTACAGGCAAGCAAGATGCCCACACAACAGGCTTAT 3180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3125 ACCTCCTGAACTGGGAACAAAGATCTACAGGCAAGCAAGATGCCCACACAACAGGCTTAT 3184

Qy    3181 TTTCTGTGAAGGAACCAACTGATCTCCCCCACCCTTGGATTAGAGTTCCTGCTCTACCTT 3240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3185 TTTCTGTGAAGGAACCAACTGATCTCCCCCACCCTTGGATTAGAGTTCCTGCTCTACCTT 3244

Qy    3241 ACCCACAGATAACACATGTTGTTTCTACTTGTAAATGTAAAGTCTTTAAAATAAACTATT 3300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3245 ACCCACAGATAACACATGTTGTTTCTACTTGTAAATGTAAAGTCTTTAAAATAAACTATT 3304

Qy    3301 ACAGATAAAAAA 3312
           ||||||||||||
Db    3305 ACAGATAAAAAA 3316
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